

re-MN

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/501,611A
Source: JFSP
Date Processed by STIC: 5/12/06

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Ne-mun

IFWP

RAW SEQUENCE LISTING

DATE: 07/14/2006

PATENT APPLICATION: US/10/501,611A

TIME: 14:48:49

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1 <110> APPLICANT: Pavan, William J.
2     Loftus, Stacie K.
3     The Government of the United States of America
4     as represented by The Secretary of the
5     Department of Health and Human Services
6 <120> TITLE OF INVENTION: Alteration of RAB38 Function to Modulate Mammalian
7     Pigmentation
8 <130> FILE REFERENCE: 015280-148100US
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/501,611A
10 <141> CURRENT FILING DATE: 2004-07-14
11 <150> PRIOR APPLICATION NUMBER: US 60/349,929
12 <151> PRIOR FILING DATE: 2002-01-18
13 <150> PRIOR APPLICATION NUMBER: WO PCT/US03/01622
14 <151> PRIOR FILING DATE: 2003-01-17
15 <160> NUMBER OF SEQ ID NOS: 28
16 <170> SOFTWARE: PatentIn Ver. 2.1
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19 <211> LENGTH: 8
20 <212> TYPE: DNA
21 <213> ORGANISM: Mus musculus
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23 <223> OTHER INFORMATION: Rab38 sequence of wildtype allele in C57Bl6/J +/+
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36     ctggktgt
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49 Asn Phe
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62 Asn Phe
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126      agctgcttgtt gatcggcgac ctgggtgtgg gcaagaccag cattatcaag cgctatgtgc 180
127      accaaaactt ctcctcgcac taccgggcca ccattggtgt ggacttcgcg ctgaagggtgc 240
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131      agtaaacgct ccctaatttgtt aagccagtgt cagtggttct gttggccaaac aaatgtgacc 480
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133      gcttcgttagg atgggttgaa acatcagccaa aggaaaacat aaacattgtat gaaggctcaa 600
134      gatgccttgtt caagcacata cttgcaaattt agtgtgacccct cctagagttt atagaaccgg 660
135      acatttgtgaa gccccatctc acatggccca aggttgtcag ctgctctggc tggccaaat 720
136      cctagaaggc tcctctgtt gcatatgaca gacagaaccc gtggccctca tgaatctgc 780
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138      gccaaagata catgactgtt tggttccctgt caaagaggaa cagcaaattttt tctttatgtt 900
139      tttccacc ccatcagcac agtgtttaca agttttttttt atattttttt gtcacaat 960
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141      taaatcaaca aaggcttctg gtcttcttgc gaaaggaaat aacagagcaa ggcagaggcc 1080
142      aagctaagtgtt tggggattttt tttttttttt gttgttctt gttcaggatcaatgtt 1140
143      ccgggtggtc tgataggtct attaaataga aaccattcat ggttagaccta aggggttgkct 1200
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145      tgccctgaat gctggttaa ctgaaaactt tatggaaaga tctgtccctt gtatgtgcct 1320
146      ttctttcagc ttcttctgac tcaagctgca ggactttctt gttatgtggaa gataattttt 1380
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150 <211> LENGTH: 291
151 <212> TYPE: DNA
  
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152 <213> ORGANISM: Homo sapiens
153 <220> FEATURE:
154 <223> OTHER INFORMATION: Rab38 exon 1 and surrounding intron sequence
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157      aggctccgca agaccccgcg gcctccagga tgcagacacc tcacaaggag caccgttaca 120
158      agctgcttgtt gatcggcgcac ctgggtgtgg gcaagaccag cattatcaag cgctatgtgc 180
159      accaaaaactt ctccctgcac taccgggcca ccattgggtt ggacttcgcg ctgaagggtgc 240
160      tccactggga cccagagacg gtggtgcgct tgcagctctg ggacattgct g 291
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163 <211> LENGTH: 281
164 <212> TYPE: DNA
165 <213> ORGANISM: Homo sapiens
166 <220> FEATURE:
167 <223> OTHER INFORMATION: Rab38 exon 2
168 <400> SEQUENCE: 11
169      gtcaagaaaat atttggaaac atgacaagag tttattaccg ggaagctatg ggggcattta 60
170      ttgtttttaa tgcaccaga ccagccacat ttgaagccgt ggcaaagtgg aaaaatgatt 120
171      tggactcaaa gttaacgctc cctaattggta agccagtgtc agtggttctg ttggccaaca 180
172      aatgtgacca agggaaaggat gtgcttatga acaatggact caagatggac cagttctgca 240
173      aggagcatgg cttcgttagga tggttgaaa catcagccaa g 281
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176 <211> LENGTH: 868
177 <212> TYPE: DNA
178 <213> ORGANISM: Homo sapiens
179 <220> FEATURE:
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184      ggttgtcagc tgctctggct gtgccaatc ctagaaggct cctctgtgg catatgacag 180
185      acagaaccccg tggccctcat gaatcgtgt tcagttttc cttattacca ttttggtaa 240
186      gcgtcaggat agggaaagcac atgtgacaaag ccaaagatac atgactgtat gttcctgtc 300
187      aaagaggaac agcaaatgtt ctttatgtgt tttcccaccc catcagcaca gtgttacaa 360
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189      gacacagaca gcccataat ttgttcctt aaatcaacaa aggctctgg tcttcttgag 480
190      aaggggaaata acagagaag gcagagggtca agctaagtgt ggggatttgc cttgccctgg 540
191      tgtgtctttg ttcaggtatc aatttgttcc cgggtggct gataggtcta ttaaatagaa 600
192      accattcatg gtagaccaa gggttgkctg ttagtggctt cttcagagtc gtgtgcacag 660
193      gcagcctggg cttttgtgt cacttgtgt gcccataatg ctggtttaac tgaaaactgt 720
194      atggaaagat ctgctccctg tatgtgcctt tctttcagct tccctgtact caagctgcag 780
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201 <213> ORGANISM: Artificial Sequence
202 <220> FEATURE:
203 <223> OTHER INFORMATION: Description of Artificial Sequence:synthetic TYRP15'T3F DNA

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210 <213> ORGANISM: Artificial Sequence
211 <220> FEATURE:
212 <223> OTHER INFORMATION: Description of Artificial Sequence:synthetic TYRP15'T7R DNA
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214 ggcgtaata cgactcaactt tagggcccag ttgcaaaatt ccagt 45
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217 <211> LENGTH: 47
218 <212> TYPE: DNA
219 <213> ORGANISM: Artificial Sequence
220 <220> FEATURE:
221 <223> OTHER INFORMATION: Description of Artificial Sequence:synthetic MLSN R T7 DNA
222 <400> SEQUENCE: 15
223 gcggtataata cgactcaactt tagggccac aaacatgtcc tacttac 47
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226 <211> LENGTH: 44
227 <212> TYPE: DNA
228 <213> ORGANISM: Artificial Sequence
229 <220> FEATURE:
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232 ggcgaaattt accctcaactt aagggaagct tccggactct ctac 44
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235 <211> LENGTH: 21
236 <212> TYPE: DNA
237 <213> ORGANISM: Artificial Sequence
238 <220> FEATURE:
239 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR
240 amplification primer Rab38 Ex1F
241 <400> SEQUENCE: 17
242 taggaaggag gattaaaccc g 21
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255 <211> LENGTH: 26
256 <212> TYPE: DNA
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 07/14/2006
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The rules require that a line not exceed 72 characters in length. This includes spaces.

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Seq#:14; Line(s) 212
Seq#:15; Line(s) 221
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VERIFICATION SUMMARY

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Input Set : N:\Crf4\Refhold\10_folder\J501611A.raw
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